

Fig. 3b

Amino acid sequence for human TNFa 8

Accession code: Swissprot P01375

- MSTESMIRDV ELAEEALPKK TGGPQGSRRC LFLSLFSFLI VAGATTLFCL -76
- -26 LHFGVIGPQR EEFPRDLSLI SPLAQA
- VRSSSRTPSD KPVAHVVANP QAEGQLQWLN RRANALLANG VELRDNQLVV 1
- PSEGLYLIYS QVLFKGQGCP STHVLLTHTI SRIAVSYQTK VNLLSAIKSP 51
- 101 CQRETPEGAE AKPWYEPIYL GGVFQLEKGD RLSAEINRPD YLDFAESGQV
- 151 YFGIIAL

Conflicting sequence F->s at position -14. Disulphide bond between Cys69-101. Signal anchor sequence -41 to -21 (underlined). Myristylation on Lys-58/-57.

Fig. 3a

	Ţ	cacaccctga	caagctgcca	ggcaggttct	cttcctctca	catactgacc
	21	cacggctcca	ccctctctcc	cctggaaagg	acaccatgag	cactgaaagc
	IOI	augaticeggg	acgtggagct	ggccgaggag	gcgctcccca	agaagacagg
	121	aaaaccccaa	ggctccaggc	ggtgcttgtt	cctcagcctc	ttctccttcc
	201	tgatcgtggc	aggcgccacc	acgetettet	gcctgctgca	ctttggagtg
	201	accygecee	agagggaaga	gtcccccagg	gacctctctc	taatcagccc
	263 20T	torggeceag	gcagtcagat	catcttctcg	aaccecgagt	gacaagcctg
	401	cageceatge	tgtagcaaac	cctcaagctg	aggggcaget	ccagtggctg
	AOT	aaccgccggg	ccaatgccct	cctggccaat	agcatagaac	toagagataa
	42T	ccadctddtd	gtgccatcag	agggcctgta	cctcatctac	tcccaggtcc
	501	tcttcaaggg	ccaaggetge	ccctccaccc	atotoctect	cacccacacc
ana a	551	atcagoogca	tegeegtete	ctaccagacc	aaggtcaacc	tectetetge
⇒ Fi	601	catcaagagc	ccctgccaga	ggagacccc	agagggggct	gaggccaagc
₩ ==	651	cctggtatga	gcccatctat	ctaagaaaaa	tettecaget	ggagaagggt
ř	701	gaccgactca	qeqetqaqat	caatcooccc	gactatetes	actttgccga
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Ų	751	gtctgggcag	gtctactttg	ggatcattgc	cctgtgagga	ggacgaacat
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=	82T	ttcagacacc	ctcaacctct	tctggctcaa	aaaqaqaatt	gggggcttag
=	Ant	ggccggaacc	caagcttaga	actttaagca	acaagaccac	cacttcgaaa
=	951	cctgggattc	aggaatgtgt	ggcctgcaca	gtgaagtget	ggcaaccact
ė						
1	1001	-	actggggcct	ccagaactca	ctggggccta	cagctttgat
	1051	•	tggaatctgg	agaccaggga	gcctttggtt	ctggccagaa
7	1101	tgctgcagga	cttgagaaga	cctcacctag	aaattgacac	aagtggacct
	1101	taggcettee	tctctccaga	tgtttccaga	cttccttgag	acacggagcc
	1201	cagecetece	catggagcca	gctccctcta	tttatgtttg	cacttgtgat
	1251	tatttattat	ttatttatta	tttatttatt	tacadatdaa	tgtatttatt
	1301	taggagacca	gogtatecto	addayccaa	tatagaegaa	gcettggctc
	1351	agacatottt	tccataaaa	cogaggetga	acaataggct	gttcccatgt
	1401	agececetaa	cctctatacc	ttcttttgat	tatettttt	aaaatattat
	1451	ctgattaaqt	totctaaaca	atgctgattt	ggtgaccaac	tgtcactcat
	1501	tgctgaggcc	tctgctcccc	agggagttgt	gtctgtaatc	ggcctactat
	1551	tcagtggcga	gaaataaagg	ttgcttagga	aagaa	
					_	

Fig. 4a

Location of inserted epitopes

Compared the compared to the c	TNF-WT
。 10. 10. 10. 10. 10. 10. 10. 10. 10. 10.	TNF2-1
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Company of the second s	TNF2-4
Windows and the second of the	TNF2-5
	TNF2-7
THE COLD STATE OF THE COLD STATE OF THE STAT	TNF30-1
	TNF30-2
A CONTRACTOR OF THE PROPERTY O	TNF30-3
policies out transfer and the contract to the	TNF30-4
TO THE PROPERTY OF THE PROPERT	TNF30-5

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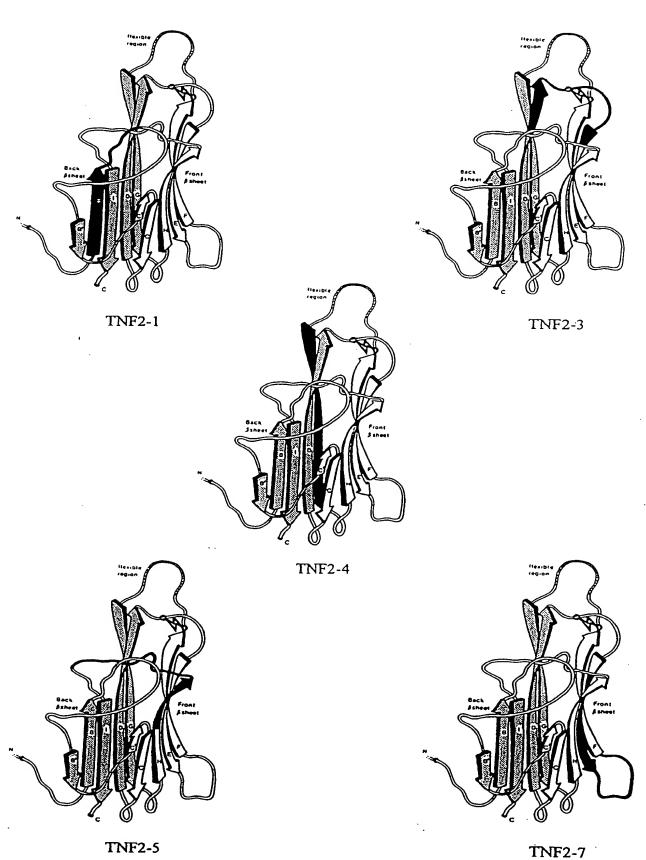
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HHH	IKA VSA			
VLLT GITE	QYIKA	IIAL		
71 STH SKFI	FWL	151 YFG:		LE
71 LFKGQGCP STHVLLTHTQYIKANSKFIGITEL-	FTVS	SGQV		 SASH
7LFKG	FNNFTVS	DFAE	IGITEL	
61 S QV 		141 D YLI		
XLIY 	ASHL	INRP	ANSK	TVSF
31	VPKVSASHLE	O1 141 151 CORETPEGAE AKPWYEPIYL GGVFQLEKGD RLSAEINRPD YLDFAESGQV YFGIIAL	-QYIKANSKF	FINNFTVSFW
51 VV P		131 GD RL		
SDNQI	NNFTVSFWLR	QLEK	KFIGITEL-	11 11 11 11 11
VELF		121 GGVF	KFIG	SASHLE
CANG		TYL	CANS	PKV
ANAL		WYE!	QYIKANS	VSFWLRVPKV
31 KRZ		111 AKE		
LOWL	YSHL!	EGAE		FNNFT
1 QAEGQLQW ITEL	VPKVSASHLE	L SRETE		
21 4P Qi	1164111	101 3P CQ		1111
HVVAI NSKF	INFTVSFWLF	SAIKS		
1 2	NNFTVSFWLI	1 VNLLS		
PSD 0 1	(4, 1	91 QTK VI		
SSRT		AVSY	NSKFIGITEL	SHLE
VRS		81 91 1 SRIAVSYQTK VNLLSAIKSP	NSK	SHL
-2-1 2-3	2-5 -2-7 -30-1 30-2 30-3 30-4		2-5 2-5 2-7 30-1	30-2 30-3 30-4 30-5
7	77	1 × 14 (വലക

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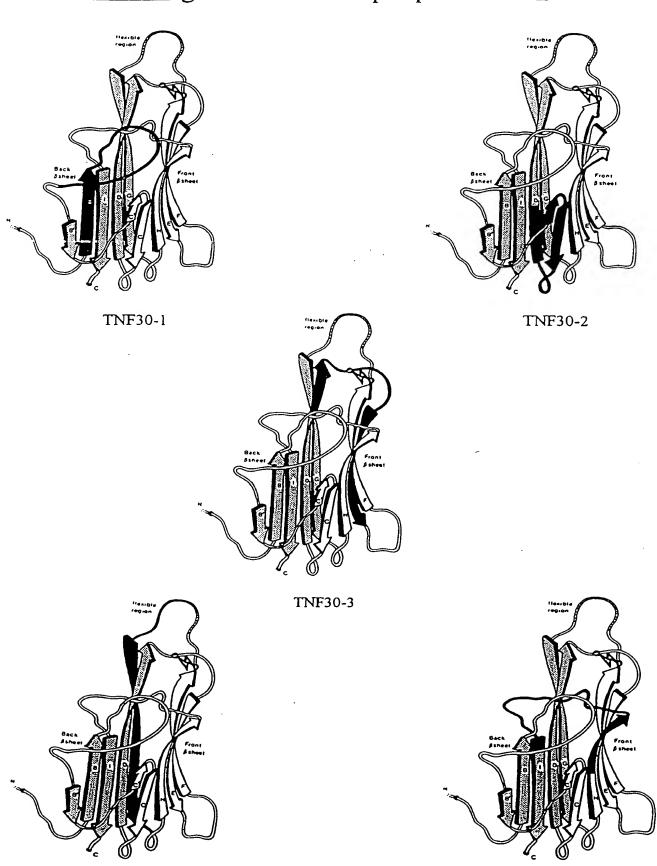
Fig. 5a

$TNF\alpha$ analogs with the P2 epitope inserted.

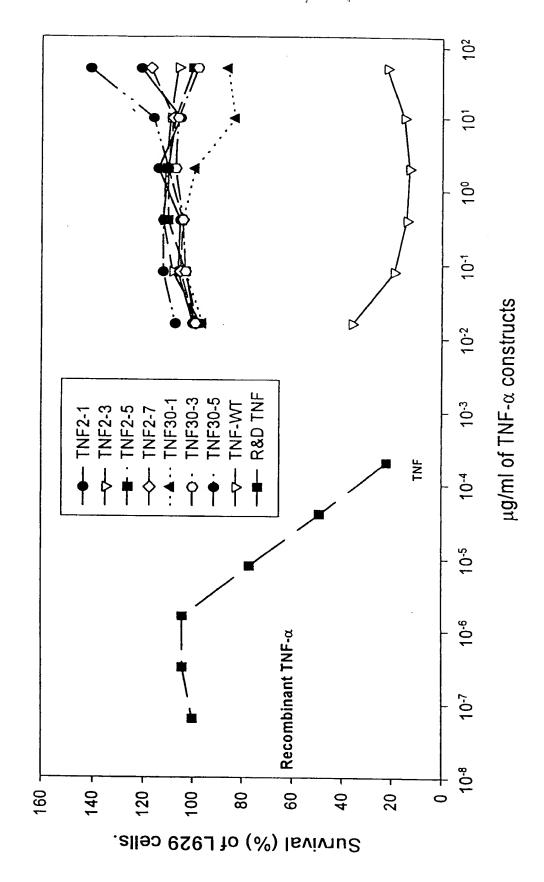


TNF30-4

TNF α analogs with the P30 epitope inserted.

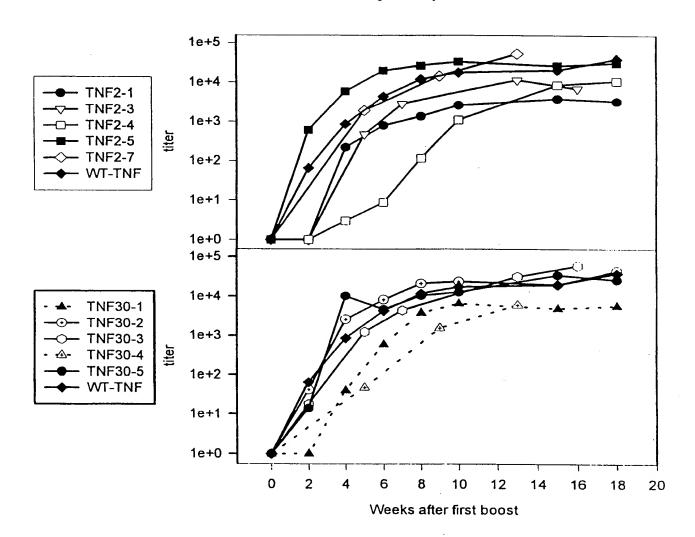


TNF30-5

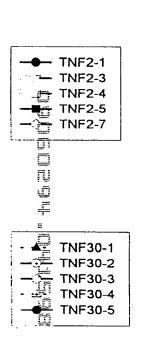


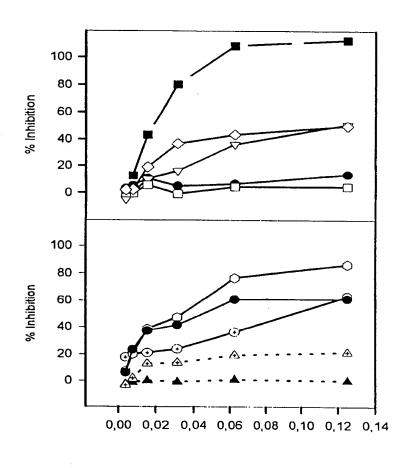
The anti human TNF α antibody responce in rabbits.

Fig. 7

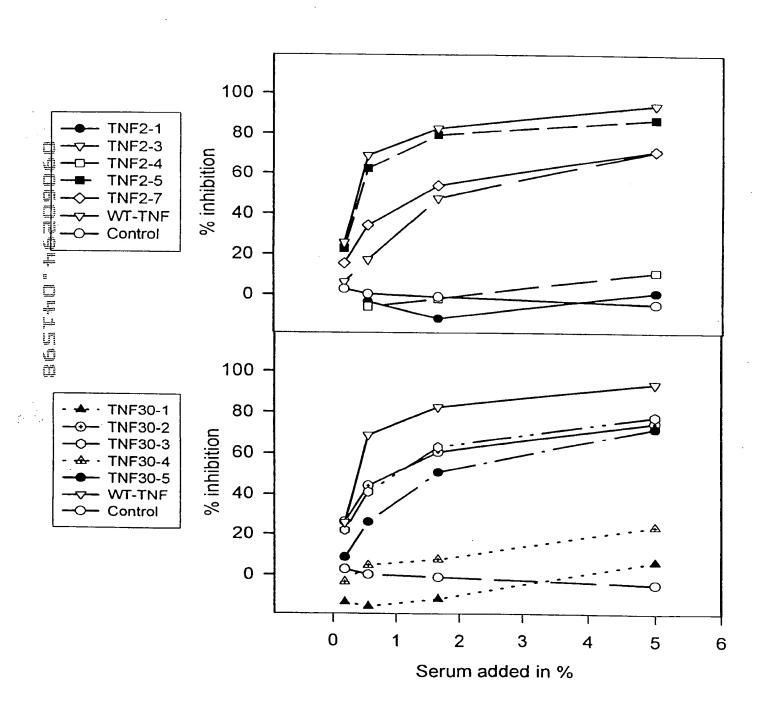


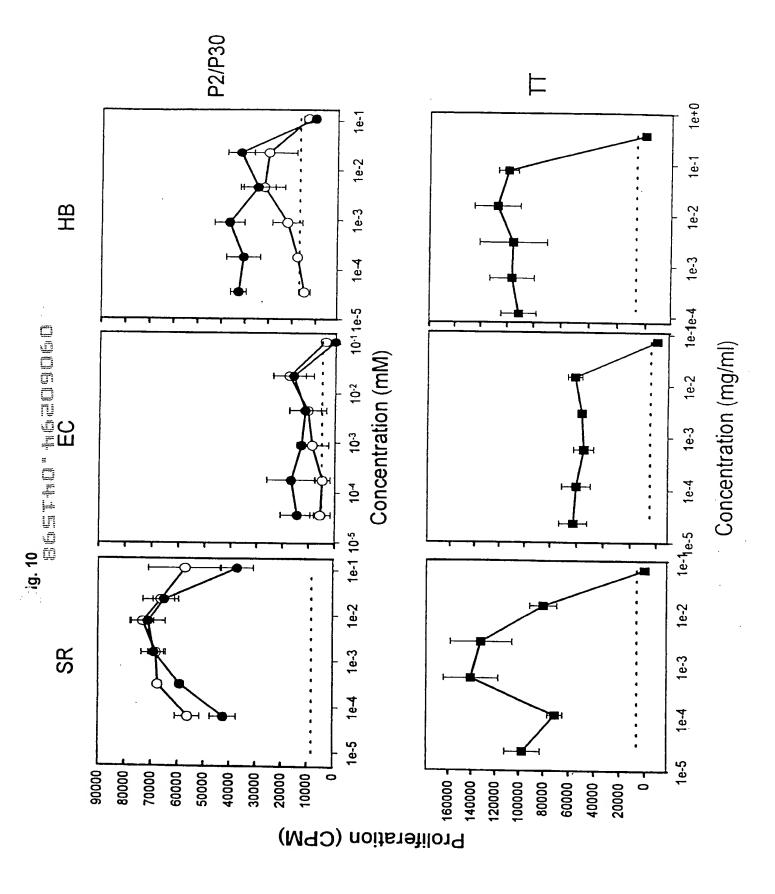
The ability of P2/P30 modified human TNF α molecules to induce neutralizing antibodies as measured in the L929 cell assay.

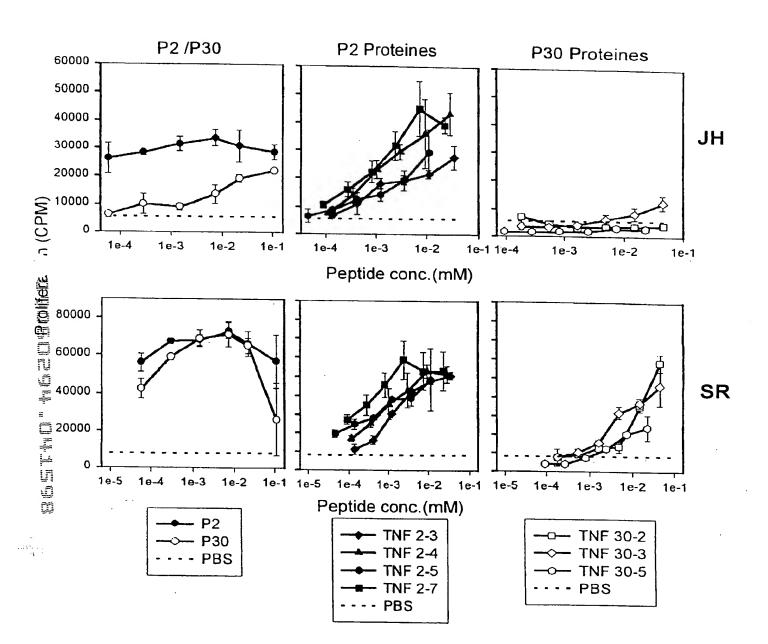




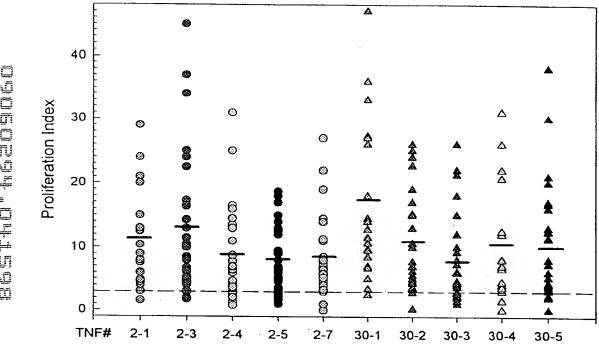
The ability of P2/P30 modified human TNF molecules to induce neutralizing antibodies as measured in the receptor assay







PBMC Assays



DOOBERT LOALES

The PBMC responce against P2 and P30 modified TNF α proteins in P2 and P30 specific responders, respectively.

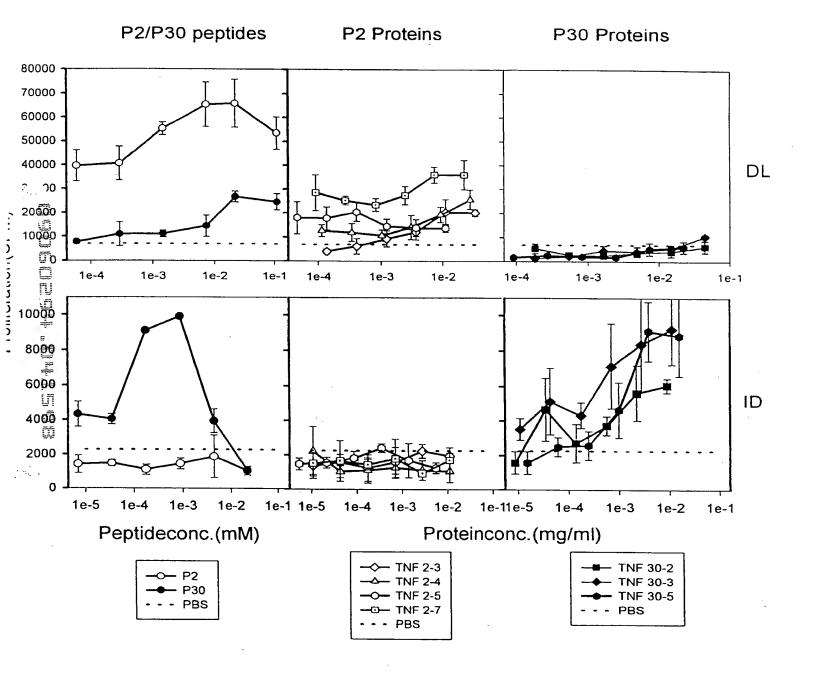


Fig. 14

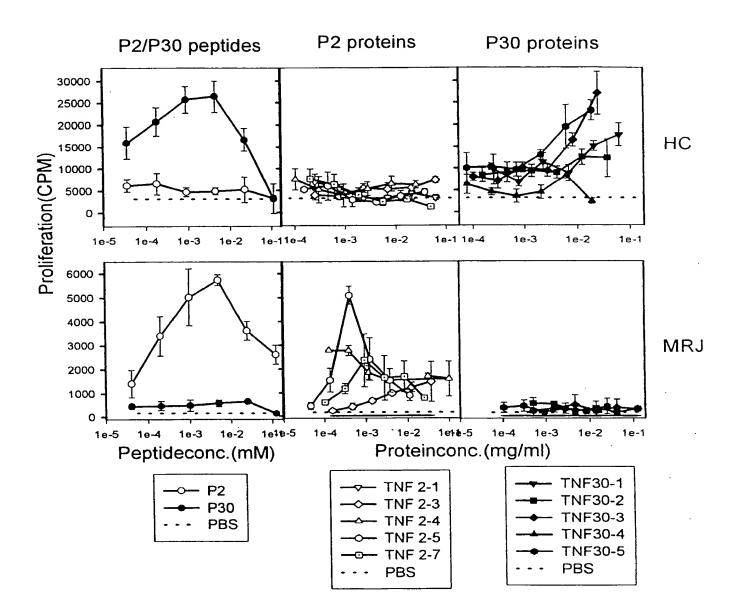
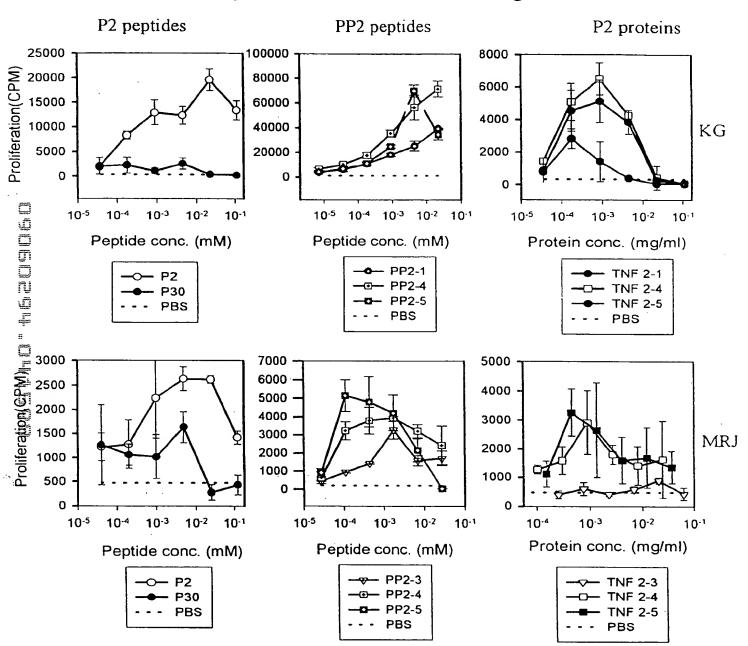
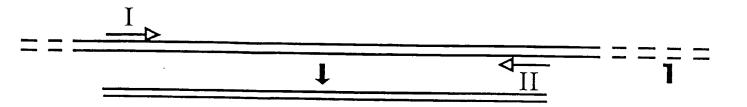


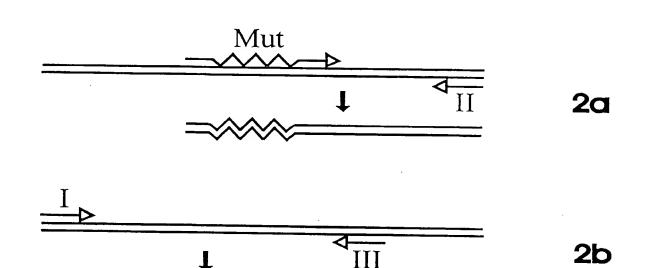
Fig. 15

The influence of flanking amino acids on the T cell recognition of P2 and P30



Mutation Strategy





Dansoa.

